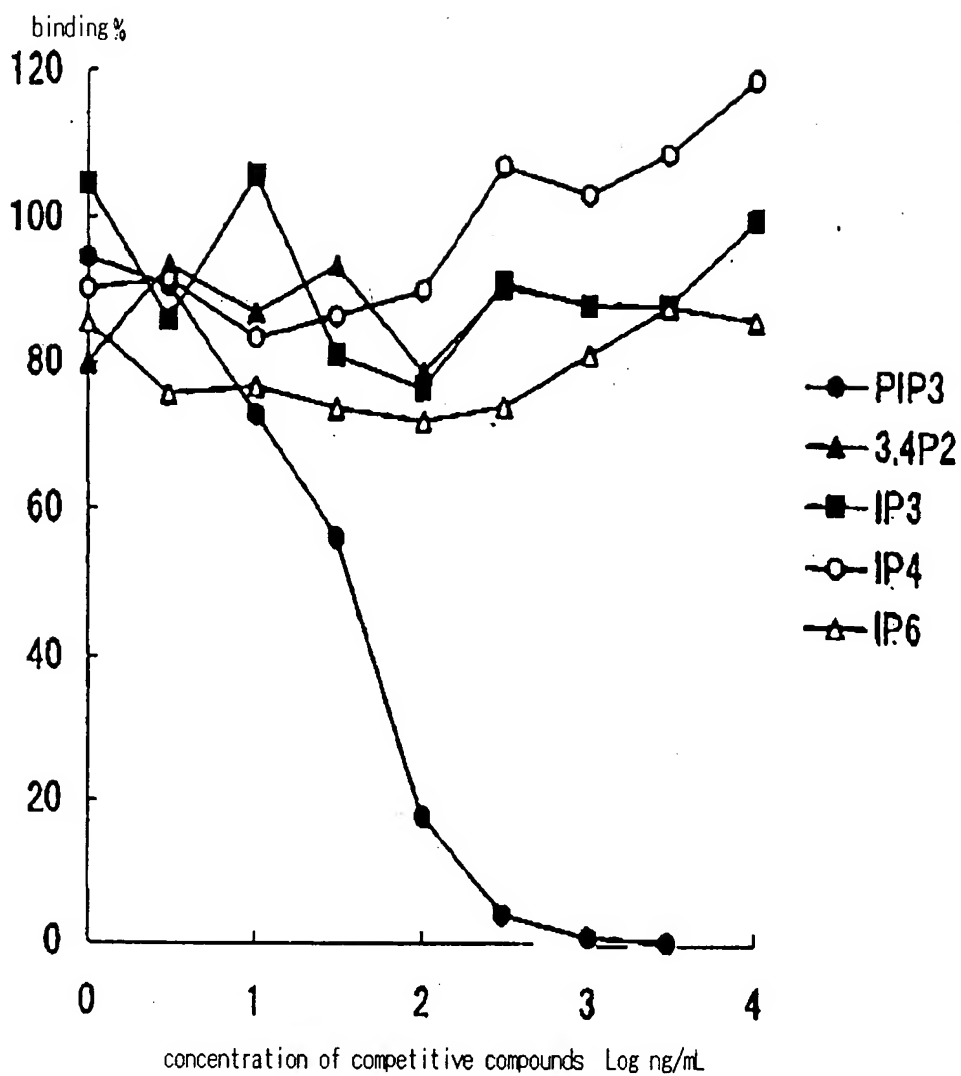


Fig.1



1000790-11301

Fig.2

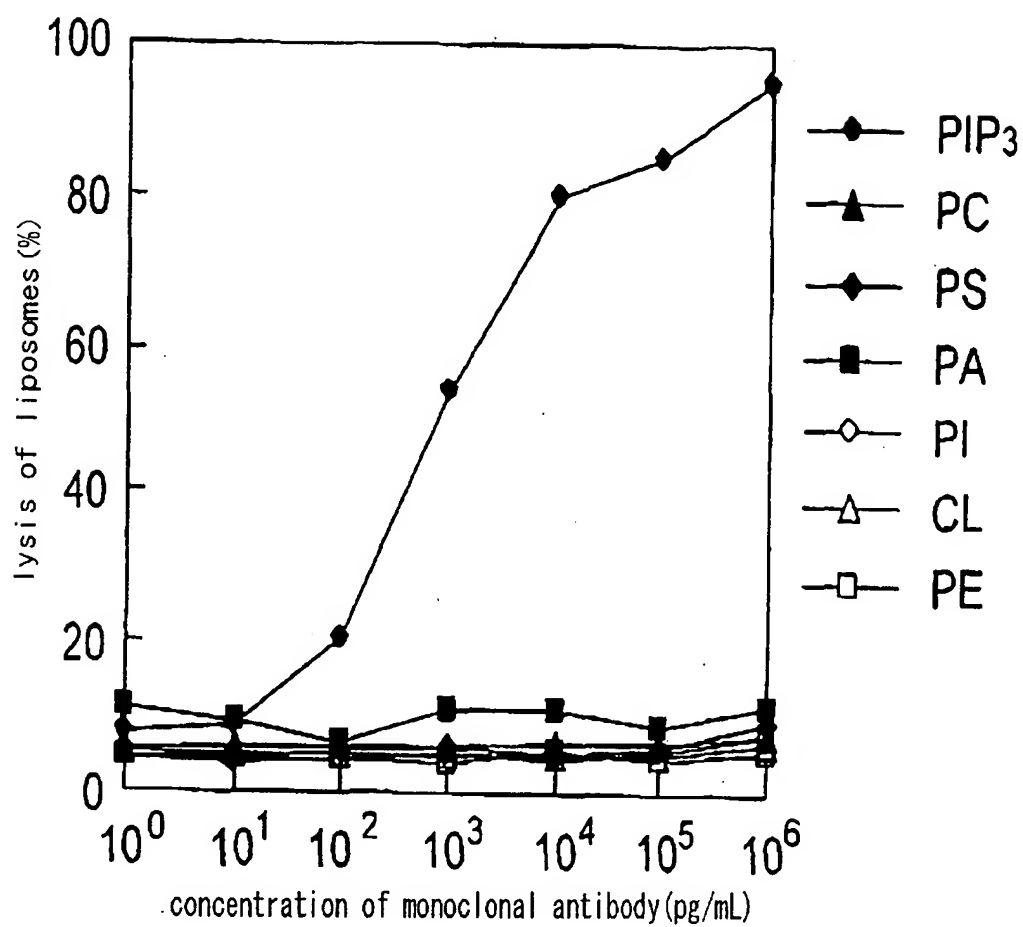


Fig.3

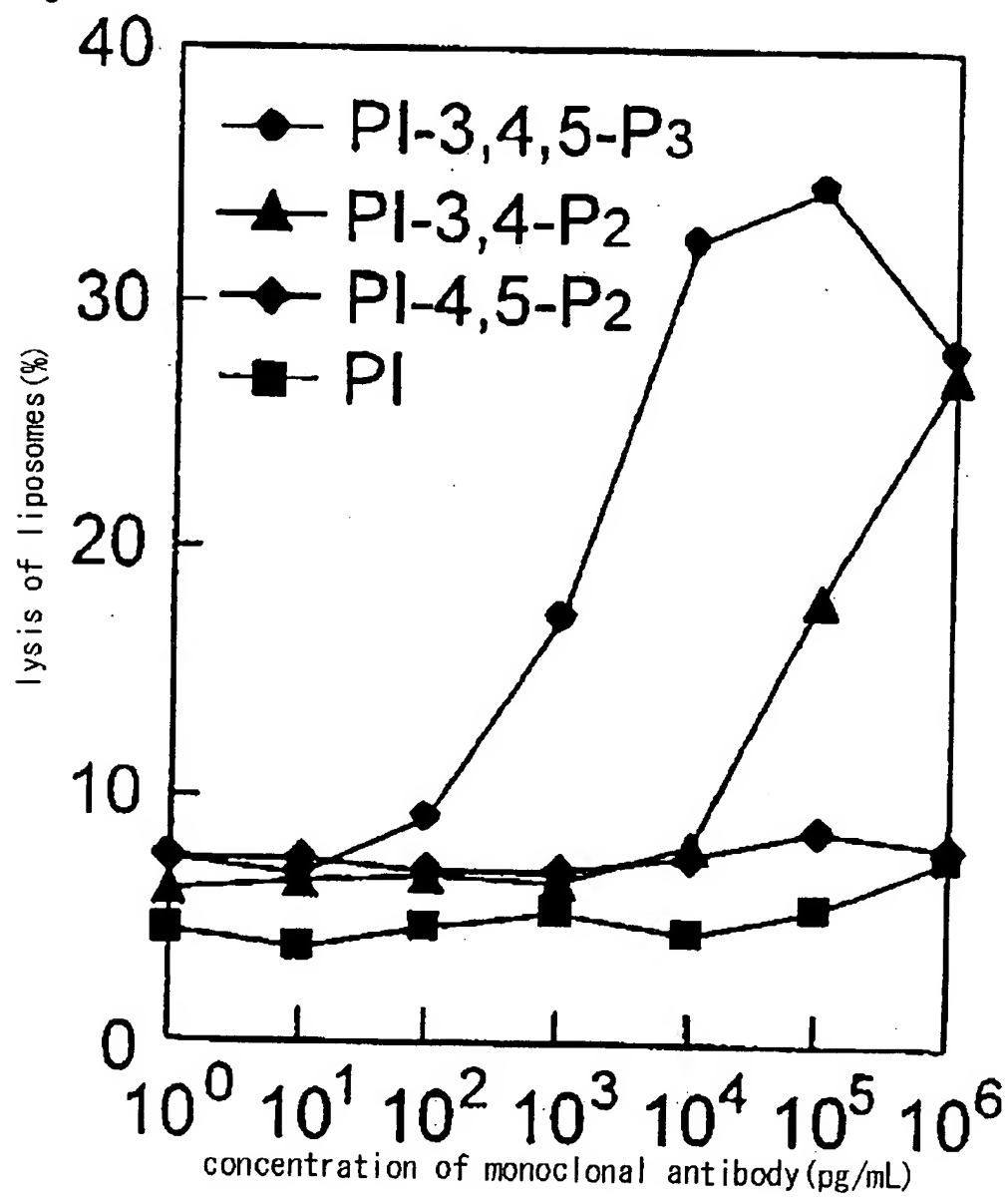
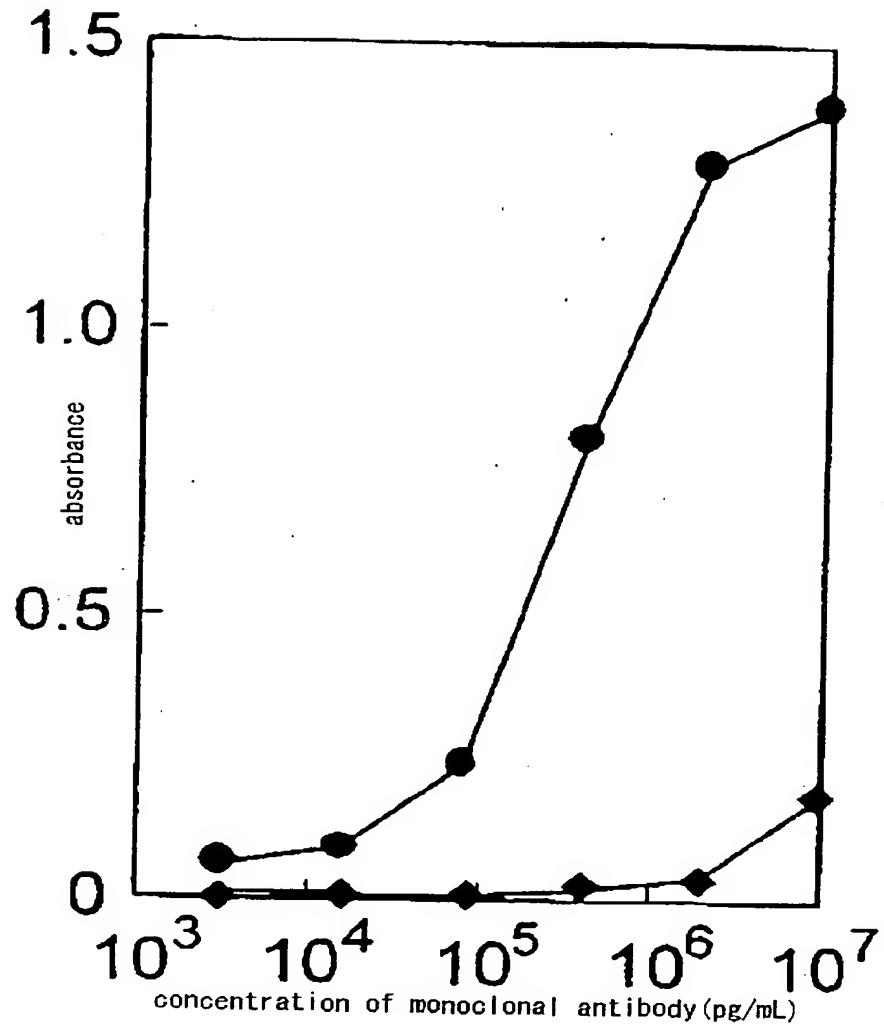
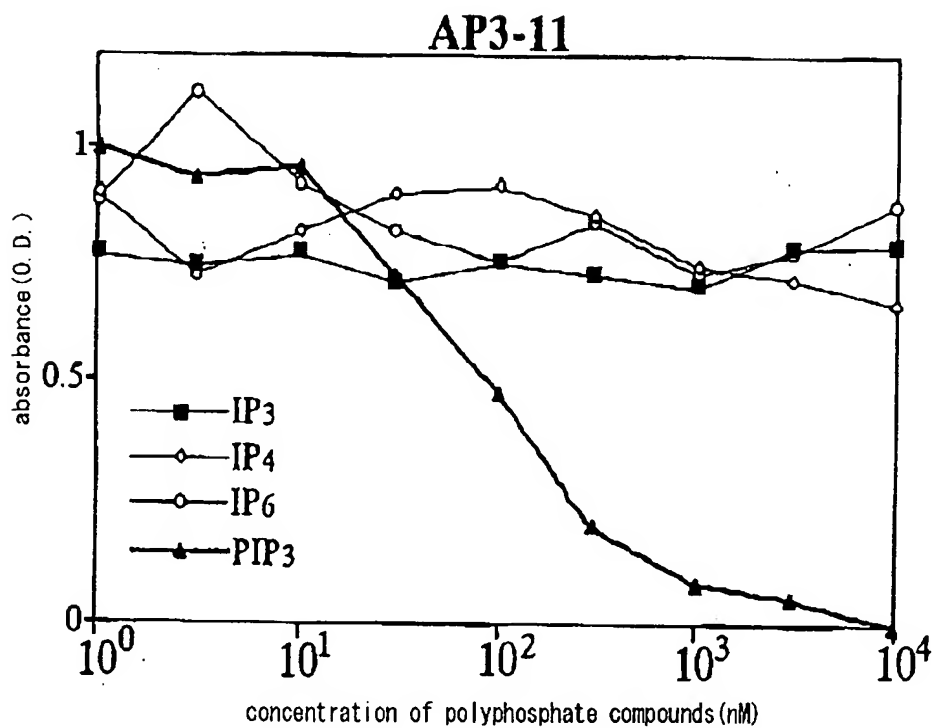
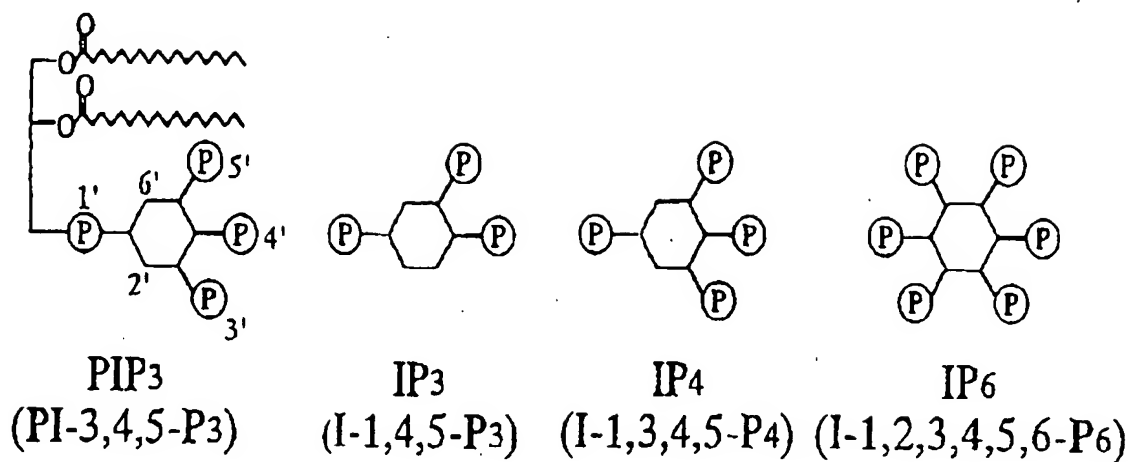


Fig.4



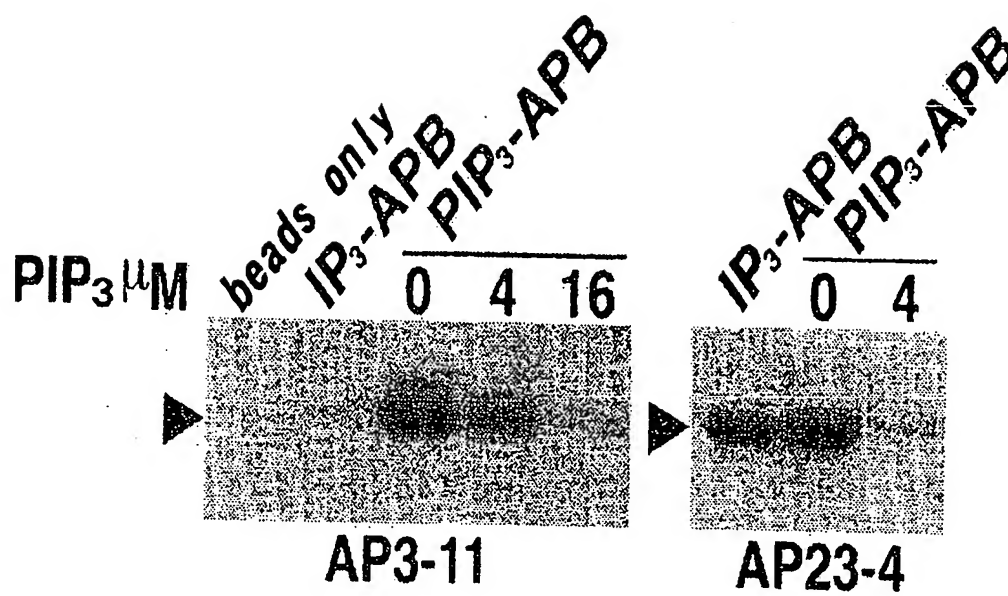
1000790-11301
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Fig.5



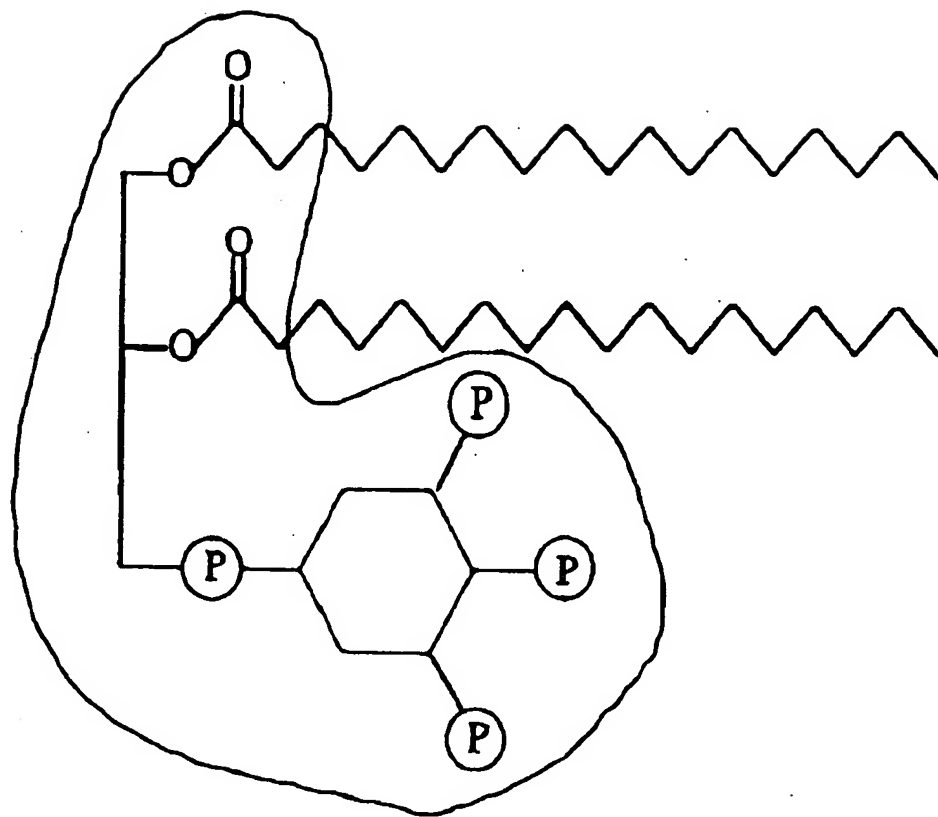
1000790-11301

Fig.6



10000790.111301

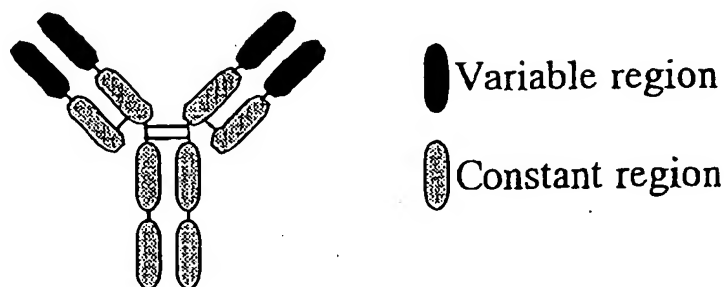
Fig.7



10007790.111301

Fig.8

Amino acid Sequence of AP3-11 variable regions



Light Chain Variable Region

DIQMTQSPASLSASVGETVTITC CDR L1
RASENIYSYLP WYQQ
 ▼ NQ GK SP QLLVY CDR L2
NAKTLTE GVPSRFSGSGSGTQFSLKIN
 SLQPEDFGSYYC CDR L3
QHHYGTPYT FGGGGTKLEIK ▼

Heavy Chain Variable Region

EVQLQQSGPEVVKT ▼ GASVKISCKAS CDR H1
GYSFTGY YMHWV
 KQSHGKSLEWIGYI CDR H2
SCYNA ATRYNQKFKGKATFTVDTS
 SSTAYMQFNSLTSEDSAVYYCAR CDR H3
SGAYYRYGVVY WGQ ▼
 GTLVTVSAATTT

FOE T T " 0642000T